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RAW SEQUENCE LISTING DATE: 08/03/2004
PATENT APPLICATION: **US/09/679,643** TIME: 14:09:20

Input Set : **N:\DA\US09679643.raw**
Output Set: **N:\CRF4\08022004\I679643.raw**

<110> APPLICANT: Foster, Timothy J.
McDevitt, Damien L.
<120> TITLE OF INVENTION: The S. aureus Fibrinogen Binding Protein Gene
<130> FILE REFERENCE: 05344.105011
<140> CURRENT APPLICATION NUMBER: US/09/679,643
<141> CURRENT FILING DATE: 2000-10-05
<150> PRIOR APPLICATION NUMBER: US/08/293,728
<151> PRIOR FILING DATE: 1994-08-22
<160> NUMBER OF SEQ ID NOS: 20
<170> SOFTWARE: PatentIn Ver. 2.0

ENTERED

<210> SEQ ID NO: 1
<211> LENGTH: 3498
<212> TYPE: DNA
<213> ORGANISM: Staphylococcus aureus
<220> FEATURE:
<221> NAME/KEY: CDS
<222> LOCATION: (301)..(3099)
<400> SEQUENCE: 1

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attcttaaaa tagcaataaa taaaatgttt gttagtaaag tattattgtg gataataaaa 180
tatcgataca aattaattgc tataatgcaa ttttagtgta taattccatt aacagagatt 240
aaatatatct taaagggtat atagttaata taaaatgact ttttaaaaag agggaataaa 300
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Ser Ala Ser Asn Glu Ser Lys Ser Asn Asp Ser Ser Ser Val Ser Ala
       50                  55                  60
gca cct aaa aca gac gac aca aac gtg agt gat act aaa aca tcg tca   540
Ala Pro Lys Thr Asp Asp Thr Asn Val Ser Asp Thr Lys Thr Ser Ser
 65                  70                  75                  80
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gaa acg aca caa tca tca tca aca aat gca act acg gaa gaa acg ccg   636
Glu Thr Thr Gln Ser Ser Ser Thr Asn Ala Thr Thr Glu Glu Thr Pro
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48              115                 120                 125
49      gca aca act caa tca agc aat aca aat gcg gag gaa tta gtg aat caa   732
50      Ala Thr Thr Gln Ser Ser Asn Thr Asn Ala Glu Glu Leu Val Asn Gln
51          130                 135                 140
52      aca agt aat gaa acg act ttt aat gat act aat aca gta tca tct gta   780
53      Thr Ser Asn Glu Thr Thr Phe Asn Asp Thr Asn Thr Val Ser Ser Val
54      145                 150                 155                 160
55      aat tca cct caa aat tct aca aat gcg gaa aat gtt tca aca acg caa   828
56      Asn Ser Pro Gln Asn Ser Thr Asn Ala Glu Asn Val Ser Thr Thr Gln
57                      165                 170                 175
58      gat act tca act gaa gca aca cct tca aac aat gaa tca gct cca cag   876
59      Asp Thr Ser Thr Glu Ala Thr Pro Ser Asn Asn Glu Ser Ala Pro Gln
60                  180                 185                 190
61      agt aca gat gca agt aat aaa gat gta gtt aat caa gcg gtt aat aca   924
62      Ser Thr Asp Ala Ser Asn Lys Asp Val Val Asn Gln Ala Val Asn Thr
63              195                 200                 205
64      agt gcg cct aga atg aga gca ttt agt tta gcg gca gta gct gca gat   972
65      Ser Ala Pro Arg Met Arg Ala Phe Ser Leu Ala Ala Val Ala Ala Asp
66          210                 215                 220
67      gca ccg gca gct ggc aca gat att acg aat cag ttg acg aat gtg aca   1020
68      Ala Pro Ala Ala Gly Thr Asp Ile Thr Asn Gln Leu Thr Asn Val Thr
69      225                 230                 235                 240
70      gtt ggt att gac tct ggt acg act gtg tat ccg cac caa gca ggt tat   1068
71      Val Gly Ile Asp Ser Gly Thr Thr Val Tyr Pro His Gln Ala Gly Tyr
72                      245                 250                 255
73      gtc aaa ctg aat tat ggt ttt tca gtg cct aat tct gct gtt aaa ggt   1116
74      Val Lys Leu Asn Tyr Gly Phe Ser Val Pro Asn Ser Ala Val Lys Gly
75                  260                 265                 270
76      gac aca ttc aaa ata act gta cct aaa gaa tta aac tta aat ggt gta   1164
77      Asp Thr Phe Lys Ile Thr Val Pro Lys Glu Leu Asn Leu Asn Gly Val
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79      act tca act gct aaa gtg cca cca att atg gct gga gat caa gta ttg   1212
80      Thr Ser Thr Ala Lys Val Pro Pro Ile Met Ala Gly Asp Gln Val Leu
81          290                 295                 300
82      gca aat ggt gta atc gat agt gat ggt aat gtt att tat aca ttt aca   1260
83      Ala Asn Gly Val Ile Asp Ser Asp Gly Asn Val Ile Tyr Thr Phe Thr
84      305                 310                 315                 320
85      gac tat gta aat act aaa gat gat gta aaa gca act ttg acc atg ccc   1308
86      Asp Tyr Val Asn Thr Lys Asp Asp Val Lys Ala Thr Leu Thr Met Pro
87                      325                 330                 335
88      gct tat att gac cct gaa aat gtt aaa aag aca ggt aat gtg aca ttg   1356
89      Ala Tyr Ile Asp Pro Glu Asn Val Lys Lys Thr Gly Asn Val Thr Leu
90                  340                 345                 350
91      gct act ggc ata ggt agt aca aca gca aac aaa aca gta tta gta gat   1404
92      Ala Thr Gly Ile Gly Ser Thr Thr Ala Asn Lys Thr Val Leu Val Asp
93              355                 360                 365
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97      gac caa atc gat aaa aca aat aat acg tat cgt cag aca att tat gtc   1500
98      Asp Gln Ile Asp Lys Thr Asn Asn Thr Tyr Arg Gln Thr Ile Tyr Val
99      385                 390                 395                 400
100      aat cca agt gga gat aac gtt att gcg ccg gtt tta aca ggt aat tta   1548
101      Asn Pro Ser Gly Asp Asn Val Ile Ala Pro Val Leu Thr Gly Asn Leu
102                      405                 410                 415
103      aaa cca aat acg gat agt aat gca tta ata gat cag caa aat aca agt   1596
104      Lys Pro Asn Thr Asp Ser Asn Ala Leu Ile Asp Gln Gln Asn Thr Ser
105                  420                 425                 430
106      att aaa gta tat aaa gta gat aat gca gct gat tta tct gaa agt tac   1644
107      Ile Lys Val Tyr Lys Val Asp Asn Ala Ala Asp Leu Ser Glu Ser Tyr
108              435                 440                 445
109      ttt gtg aat cca gaa aac ttt gag gat gtc act aat agt gtg aat att   1692
110      Phe Val Asn Pro Glu Asn Phe Glu Asp Val Thr Asn Ser Val Asn Ile
111          450                 455                 460
112      aca ttc cca aat cca aat caa tat aaa gta gag ttt aat acg cct gat   1740
113      Thr Phe Pro Asn Pro Asn Gln Tyr Lys Val Glu Phe Asn Thr Pro Asp
114      465                 470                 475                 480
115      gat caa att aca aca ccg tat ata gta gtt gtt aat ggt cat att gat   1788
116      Asp Gln Ile Thr Thr Pro Tyr Ile Val Val Val Asn Gly His Ile Asp
117                      485                 490                 495
118      ccg aat agc aaa ggt gat tta gct tta cgt tca act tta tat ggg tat   1836
119      Pro Asn Ser Lys Gly Asp Leu Ala Leu Arg Ser Thr Leu Tyr Gly Tyr
120                  500                 505                 510
121      aac tcg aat ata att tgg cgc tct atg tca tgg gac aac gaa gta gca   1884
122      Asn Ser Asn Ile Ile Trp Arg Ser Met Ser Trp Asp Asn Glu Val Ala
123              515                 520                 525
124      ttt aat aac gga tca ggt tct ggt gac ggt atc gat aaa cca gtt gtt   1932
125      Phe Asn Asn Gly Ser Gly Ser Gly Asp Gly Ile Asp Lys Pro Val Val
126          530                 535                 540
127      cct gaa caa cct gat gag cct ggt gaa att gaa cca att cca gag gat   1980
128      Pro Glu Gln Pro Asp Glu Pro Gly Glu Ile Glu Pro Ile Pro Glu Asp
129      545                 550                 555                 560
130      tca gat tct gac cca ggt tca gat tct ggc agc gat tct aat tca gat   2028
131      Ser Asp Ser Asp Pro Gly Ser Asp Ser Gly Ser Asp Ser Asn Ser Asp
132                      565                 570                 575
133      agc ggt tca gat tcg ggt agt gat tct aca tca gat agt ggt tca gat   2076
134      Ser Gly Ser Asp Ser Gly Ser Asp Ser Thr Ser Asp Ser Gly Ser Asp
135                  580                 585                 590
136      tca gcg agt gat tca gat tca gca agt gat tca gac tca gcg agt gat   2124
137      Ser Ala Ser Asp Ser Asp Ser Ala Ser Asp Ser Asp Ser Ala Ser Asp
138              595                 600                 605
139      tca gat tca gca agc gat tcc gac tca gcg agc gat tcc gac tca gac   2172
140      Ser Asp Ser Ala Ser Asp Ser Asp Ser Ala Ser Asp Ser Asp Ser Asp
141          610                 615                 620
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146       Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp
147                       645                 650                 655
148       tca gat tca gat agc gat tca gat tca gat agc gat tca gat tcc gac   2316
149       Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp
150                 660                 665                 670
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152       Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp
153             675                 680                 685
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155       Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp
156           690                 695                 700
157       tcc gac tca gat agc gac tca gat tca gac agc gat tca gat tca gac   2460
158       Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp
159       705                 710                 715                 720
160       agc gat tca gat tca gat agc gat tca gat tcc gac agt gac tca gat   2508
161       Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp
162                       725                 730                 735
163       tcc gac agt gac tcg gat tca gat agc gat tca gat tcc gac agt gac   2556
164       Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp
165                 740                 745                 750
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168             755                 760                 765
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171           770                 775                 780
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173       Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp
174       785                 790                 795                 800
175       tcg gac tca gat agc gat tca gaa tca gac agc gat tca gaa tca gac   2748
176       Ser Asp Ser Asp Ser Asp Ser Glu Ser Asp Ser Asp Ser Glu Ser Asp
177                       805                 810                 815
178       agc gat tca gat tca gac agc gac tca gac agt gac tca gat tca gat   2796
179       Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp Ser Asp
180                 820                 825                 830
181       agt gac tcg gat tca gcg agt gat tca gac tca ggt agt gac tcc gat   2844
182       Ser Asp Ser Asp Ser Ala Ser Asp Ser Asp Ser Gly Ser Asp Ser Asp
183             835                 840                 845
184       tca tca agt gat tcc gac tca gaa agt gat tca aat agc gat tcc gag   2892
185       Ser Ser Ser Asp Ser Asp Ser Glu Ser Asp Ser Asn Ser Asp Ser Glu
186           850                 855                 860
187       tca ggt tct aac aat aat gta gtt ccg cct aat tca cct aaa aat ggt   2940
188       Ser Gly Ser Asn Asn Asn Val Val Pro Pro Asn Ser Pro Lys Asn Gly
189       865                 870                 875                 880
190       act aat gct tct aat aaa aat gag gct aaa gat agt aaa gaa cca tta   2988
191       Thr Asn Ala Ser Asn Lys Asn Glu Ala Lys Asp Ser Lys Glu Pro Leu
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tta tta gca tca ata ggt tca tta cta ctt ttc aga aga aaa aaa gaa   3084
Leu Leu Ala Ser Ile Gly Ser Leu Leu Leu Phe Arg Arg Lys Lys Glu
        915                 920                 925
aat aaa gat aag aaa taagtaataa tgatattaaa ttaatcatat gattcatgaa   3139
Asn Lys Asp Lys Lys
    930
gaagccacct taaaaggtgc ttcttttact tggattttcc aaatatattg tttgaatata 3199
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cagttgaaaa ataaattgct agttttatca ttgggagcat tatgtgtatc acaaatttgg 3379
gaaagtaatc gtgcgagtgc agtggtttct ggggagaaga atccatatgt atctgagtcg 3439
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        35                  40                  45
Ser Ala Ser Asn Glu Ser Lys Ser Asn Asp Ser Ser Ser Val Ser Ala
    50                  55                  60
Ala Pro Lys Thr Asp Asp Thr Asn Val Ser Asp Thr Lys Thr Ser Ser
65                  70                  75                  80
Asn Thr Asn Asn Gly Glu Thr Ser Val Ala Gln Asn Pro Ala Gln Gln
                85                  90                  95
Glu Thr Thr Gln Ser Ser Ser Thr Asn Ala Thr Thr Glu Glu Thr Pro
            100                 105                 110
Val Thr Gly Glu Ala Thr Thr Thr Thr Thr Asn Gln Ala Asn Thr Pro
        115                 120                 125
Ala Thr Thr Gln Ser Ser Asn Thr Asn Ala Glu Glu Leu Val Asn Gln
    130                 135                 140
Thr Ser Asn Glu Thr Thr Phe Asn Asp Thr Asn Thr Val Ser Ser Val
145                 150                 155                 160
Asn Ser Pro Gln Asn Ser Thr Asn Ala Glu Asn Val Ser Thr Thr Gln
                165                 170                 175
Asp Thr Ser Thr Glu Ala Thr Pro Ser Asn Asn Glu Ser Ala Pro Gln
            180                 185                 190
Ser Thr Asp Ala Ser Asn Lys Asp Val Val Asn Gln Ala Val Asn Thr
        195                 200                 205
Ser Ala Pro Arg Met Arg Ala Phe Ser Leu Ala Ala Val Ala Ala Asp
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